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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/667,141	09/18/2003	Mario H. Skiadopoulos	1173-1034PUS2 7197	
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Birch, Stewar	t, Kolasch & Birch, LLP		BOESEN, A	GNIESZKA
8110 Gatehous P.O. Box 747	e Rd, Suite 500 East		ART UNIT	PAPER NUMBER
	VA 22040-0747		1648	
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Please find below and/or attached an Office communication concerning this application or proceeding.

	Application No.	Applicant(s)
	10/667,141	SKIADOPOULOS ET AL.
Office Action Summary	Examiner	Art Unit
	Agnieszka Boesen	1648
The MAILING DATE of this communication app	ears on the cover sheet with the c	orrespondence address
Period for Reply	/ 10 0 = T T	0) 00 THDT/ (00) DAY(0
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period w - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 16(a). In no event, however, may a reply be tirr rill apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	I. lely filed the mailing date of this communication. C (35 U.S.C. § 133).
Status		
1) Responsive to communication(s) filed on 27 Ap	<u>oril 2006</u> .	
2a) This action is FINAL . 2b) ⊠ This	action is non-final.	
3) Since this application is in condition for allowan		
closed in accordance with the practice under E	x parte Quayle, 1935 C.D. 11, 45	i3 O.G. 213.
Disposition of Claims		
4) Claim(s) <u>1-129,183,232,255 and 278</u> is/are per	nding in the application.	
4a) Of the above claim(s) is/are withdraw	vn from consideration.	
5) Claim(s) is/are allowed.		
6) Claim(s) is/are rejected.		
7) Claim(s) is/are objected to.		
8) Claim(s) <u>1-129,183,232,255 and 278</u> are subje	ct to restriction and/or election re	quirement.
Application Papers		
9)☐ The specification is objected to by the Examine	r.	
10) The drawing(s) filed on is/are: a) acce	epted or b) objected to by the f	Examiner.
Applicant may not request that any objection to the	drawing(s) be held in abeyance. See	e 37 CFR 1.85(a).
Replacement drawing sheet(s) including the correcti		
11) The oath or declaration is objected to by the Ex	aminer. Note the attached Office	Action or form PTO-152.
Priority under 35 U.S.C. § 119		
12) Acknowledgment is made of a claim for foreign	priority under 35 U.S.C. § 119(a)	-(d) or (f).
a) ☐ All b) ☐ Some * c) ☐ None of:		
 Certified copies of the priority documents 		
2. Certified copies of the priority documents	• • • • • • • • • • • • • • • • • • • •	
3. Copies of the certified copies of the prior		ed in this National Stage
application from the International Bureau * See the attached detailed Office action for a list		od.
See the attached detailed Office action for a list	or the certified copies not receive	·
Attachment(s)	_	
Notice of References Cited (PTO-892) Notice of Draftsperson's Patent Drawing Review (PTO-948)	4) Interview Summary Paper No(s)/Mail Da	
3) Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08) Paper No(s)/Mail Date		ratent Application (PTO-152)

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DETAILED ACTION

The Examiner and/or Art Unit location of your application in the PTO has changed. To aid in correlating any papers for this application, all further correspondence regarding this application should be directed to Examiner Agnieszka Boesen Group Art Unit 1648.

Applicants called on April 27, 2006 to question the Election/Restriction requirement mailed December 30, 2005. Applicants noted that in other cases the inventions are separated into products, methods of using and methods of making. In this instance the prior examiner had grouped methods of using and methods of making into the same group, but separated the polynucleotides as "partial" or "complete." Applicants' demanded that the Office issue a new Election/Restriction requirement, separating the products from the methods of making and using the compostions. Upon review and reconsideration the Election/Restriction requirement mailed mailed December 30, 2005 is herby vacated and the following new Election/Restriction is issued.

Restriction to one of the following inventions is required under 35 U.S.C. 121:

Group #	Linking claims	Claims	
1	67, 70	71, 72, 129	A composition comprising a "complete" HPIV2 polyhexameric genome in a particle. The genome must possess N, P and L protein The composition is isolated, infectious, self-replicating. The "recombinant" requirement is read as a process (product-by-process) limitation, because it does not add to the structure of the composition, classified in class 424, subclass 211.1.
2	67, 70	72, 83, 85, 86, 129	A composition comprising a "partial" HPIV2 polyhexameric genome in a particle. The genome must possess N, P and L protein The composition is isolated, infectious, self-replicating. The composition is isolated, infectious, self-replicating.

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			The "recombinant" requirement is read as a process (product- by-process) limitation, because it does not add to the structure
			of the composition, classified in class 424, subclass 211.1.
3	67, 73	74, 75, 76,	Attenuating mutation HPIV3 JS cp45
		255, 278	Target at 948 and/or 1566 HPIV2 L protein, classified in class
			424, subclass 211.1.
4	67, 73	74, 77, 78,	Attenuating mutation from RSV.
		82	Target at Phe 460 in HPIV2 L, classified in class 424, subclass
			211.1.
5	67, 73	74, 79, 80	Attenuating mutation from BPIV3
			Target at Ser 1724 in the HPIV2 L protein is a substitution,
			classified in class 424, subclass 211.1.
6	67, 73	74, 79, 80,	Attenuating mutation from BPIV3
		81	Target at Ser 1724-1725 in the HPIV2 L protein is a deletion,
			classified in class 424, subclass 211.1.
7	67, 83	84, 85, 86,	Nucleotide modification resulting in a phenotypic change. A
	:	124, 125-128	nucleotide modification can be an <u>insertion</u> , <u>deletion or</u>
			substitution. Resulting in the following phenotypic change:
			Attenuation
			Temperature sensitivity
			Cold-adaptation
			Plaque size
			Host-range restriction
			Change in immunogenicity, classified in class 424, subclass 211.1
8	67.02	83, 87, 88	Nucleotide modification (insertion) to encode a non-PIV
0	67, 83	03, 07, 00	molecule where the molecule is a cytokine , classified in class
			424, subclass 211.1.
9	67, 83	87	Nucleotide modification (insertion) to encode a non-PIV
	07,05	"	molecule where the molecule is a T-helper epitope , classified
			in class 424, subclass 211.1.
10	67, 83	87	Nucleotide modification (insertion) to encode a non-PIV
	07,00		molecule where the molecule is a restriction site marker.
11	67, 83	87,	Nucleotide modification (insertion) to encode a non-PIV
	, , , , ,	,	molecule where the molecule is a protein of a microbial
			(bacteria or virus) pathogen, classified in class 424, subclass
			211.1
12	67, 89	90, 94, 95,	HPIV2 chimera with a heterologous gene encoding antigenic
		98, 99, 100,	determinant. The new gene segment is added (supernumerary)
		101, 102,	in addition to the HPIV2 genes. The supernumerary
		103, 104	heterologous gene is HPIV1 HN, classified in class 424,
			subclass 211.1.
13	67, 89	100, 104	HPIV2 chimera with a heterologous gene encoding antigenic
			determinant. The new gene segment is added (supernumerary)

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			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV1 F, classified in class 424, subclass
			211.1.
14	67, 89	90, 94, 95	HPIV2 chimera with a heterologous gene encoding antigenic
	İ		determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV2 F , classified in class 424, subclass
			211.1.
15	67, 89	90, 94, 95,	HPIV2 chimera with a heterologous gene encoding antigenic
		100, 104	determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV3 HN, classified in class 424,
			subclass 211.1.
16	67, 89	90, 94, 95	HPIV2 chimera with a heterologous gene encoding antigenic
			determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV3 F, classified in class 424, subclass
			211.1.
17	67, 89	68, 92	HPIV2 chimera with at least one different HPIV N, P or L
			protein.
18	67, 89	68, 89, 120,	HPIV2 /BPIV3 chimera with at least one different BPIV3 N, P
		121, 122,	or L protein, classified in class 424, subclass 211.1.
		123	
19	67, 89	90, 94, 95	HPIV2 chimera with a heterologous gene encoding antigenic
			determinant with an added gene sequence. The supernumerary
			heterologous gene is measles virus HA, classified in class 424,
			subclass 211.1.
20	67, 89	91	HPIV2 chimera with a heterologous gene includes a regulatory
			element, classified in class 424, subclass 211.1.
21	67, 89	93, 96	HPIV2 chimera with a heterologous measles virus pathogen
			sequence, classified in class 424, subclass 212.1.
22	67, 89	93, 96, 97	HPIV2 chimera with a heterologous RSV subgroup A
		ļ	pathogen sequence, classified in class 424, subclass 211.1.
23	67, 89	93, 96, 97	HPIV2 chimera with a heterologous RSV subgroup B virus
<u> </u>		100.06	pathogen sequence, classified in class 424, subclass 207.1.
24	67, 89	93, 96	HPIV2 chimera with a heterologous mumps virus pathogen
	(5.00	100.06	sequence, classified in class 424, subclass 211.1.
25	67, 89	93, 96	HPIV2 chimera with a heterologous human papilloma virus
26	(7, 00	02.06	pathogen sequence, classified in class 424, subclass 211.1.
26	67, 89	93, 96	HPIV2 chimera with a heterologous HIV type 1 virus
07	(7.00	02.06	pathogen sequence, classified in class 424, subclass 208.1
27	67, 89	93,96	HPIV2 chimera with a heterologous HIV type 2 virus
20	(7.00	02.06	pathogen sequence, classified in class 424, subclass 208.1.
28	67, 89	93, 96	HPIV2 chimera with a heterologous HSV virus pathogen

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			sequence, classified in class 424, subclass 211.1.
29	67, 89	93, 96	HPIV2 chimera with a heterologous cytomegalovirus
			pathogen sequence, classified in class 424, subclass 211.1.
30	67, 89	93, 96	HPIV2 chimera with a heterologous rabies virus pathogen
			sequence, classified in class 424, subclass 211.1.
31	67, 89	93, 96	HPIV2 chimera with a heterologous human
			metapneumovirus pathogen sequence, classified in class 424,
			subclass 211.1.
32	67, 89	93, 96	HPIV2 chimera with a heterologous Epstein Barr virus
			pathogen sequence, classified in class 424, subclass 211.1.
33	67, 89	93, 96	HPIV2 chimera with a heterologous filovirus pathogen
			sequence, classified in class 424, subclass 211.1.
34	67, 89	93, 96	HPIV2 chimera with a heterologous bunyavirus pathogen
			sequence, classified in class 424, subclass 211.1.
35	67, 89	93, 96	HPIV2 chimera with a heterologous flavivirus pathogen
			sequence, classified in class 424, subclass 211.1.
36	67, 89	93, 96	HPIV2 chimera with a heterologous alphavirus pathogen
			sequence, classified in class 424, subclass 211.1.
37	67, 89	93, 96	HPIV2 chimera with a heterologous influenza virus pathogen
			sequence, classified in class 424, subclass 211.1.
38	67, 89	105, 106,	HPIV2 chimera with an attenuating mutation HPIV3 JS cp45
		107	Target at 948 and/or 1566 HPIV2 L protein, classified in class
			424, subclass 211.1
39	67, 89	105, 108,	HPIV2 chimera with an attenuating mutation from RSV with a
		109	substitution at corresponding target position Phe 460 in
			HPIV2 L protein, classified in class 424, subclass 211.1.
40	67, 89	105, 111,	HPIV2 chimera with an additional attenuating mutation from
		123	BPIV3. Here the target at Ser 1724 in the HPIV2 L protein is a
			substitution, classified in class 424, subclass 211.1.
41	67, 89	105, 111,	HPIV2 chimera with an additional attenuating mutation from
		112	BPIV3. Here the target at Ser 1724 in the HPIV2 L protein is a
			deletion, classified in class 424, subclass 211.1.
42	67, 89	105, 114,	HPIV2 / HPIV1 chimera that comprises an additional
		115,	attenuating mutation, classified in class 424, subclass 211.1.
43	67, 89	105, 113,	HPIV2 chimera that comprises an additional attenuating
		116, 117,	mutation (insertion deletion, substitution) resulting in any one
		118	of the following phenotypic changes:
			Attenuation
			Temperature sensitivity
			Cold-adaptation
			Plaque size
			Host-range restriction
			Change in immunogenicity
		<u> </u>	Here the attenuation is a partial or complete deletion of HPIV2

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			V, classified in class 424, subclass 211.1.
44	67, 89	116, 119	HPIV2 chimera that comprises an additional attenuating mutation (insertion) resulting in a phenotypic change where the inserted non-PIV molecule is a cytokine , classified in class 424, subclass 211.1.
45	67, 89	116, 119	HPIV2 chimera that comprises an additional attenuating mutation (insertion) resulting in a phenotypic change where the inserted non-PIV molecule is a T-helper epitope , classified in class 424, subclass 211.1.
46	67, 89	116, 119	HPIV2 chimera that comprises an additional attenuating mutation (insertion) resulting in a phenotypic change where the inserted non-PIV molecule is a restriction site marker , classified in class 424, subclass 211.1.
47	67, 89	116, 119	HPIV2 chimera that comprises an attenuating mutation (insertion) resulting in a phenotypic change where the inserted non-PIV molecule is a protein of a microbial pathogen (bacteria or virus), classified in class 424, subclass 277.1.
48	1	2, 3, 4, 5,	A method of making a composition comprising a "complete" HPIV2 polyhexameric genome in a particle. The genome must possess N, P and L protein The composition is isolated, infectious, self-replicating. The "recombinant" requirement is read as a process (product-by-process) limitation, because it does not add to the structure of the composition, classified in class 435, subclass 5.
49	1	2, 3, 4,	A method of making a composition comprising a "partial" HPIV2 polyhexameric genome in a particle. The genome must possess N, P and L protein The composition is isolated, infectious, self-replicating. The composition is isolated, infectious, self-replicating. The "recombinant" requirement is read as a process (product-by-process) limitation, because it does not add to the structure of the composition, classified in class 435, subclass 5.
50	1, 9	11, 12, 13, 14	A method of making a composition comprising an attenuating mutation HPIV3 JS cp45 Target at 948 and/or 1566 HPIV2 L protein, classified in class 435, subclass 5.
51	1, 9	15, 16	A method of making a composition comprising an attenuating mutation from RSV. Target at Phe 460 in HPIV2 L, classified in class 435, subclass 5.
52	1,9	17, 18	A method of making a composition comprising an attenuating mutation from BPIV3 Target at Ser 1724 in the HPIV2 L protein is a substitution, classified in class 435, subclass 5.

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53	1, 9	17, 18, 19	A method of making a composition comprising an attenuating
			mutation from BPIV3
			Target at Ser 1724-1725 in the HPIV2 L protein is a deletion,
			classified in class 435, subclass 5.
54	1, 21	8, 20, 21, 22,	A method of making a composition comprising a nucleotide
	i	23, 24, 25	modification resulting in a phenotypic change. A nucleotide
			modification can be an <u>insertion</u> , <u>deletion or substitution</u> .
			Resulting in the following phenotypic change:
			Attenuation
			Temperature sensitivity
			Cold-adaptation
			Plaque size
			Host-range restriction
			Change in immunogenicity, classified in class 435,
			subclass 5.
55	1, 21,	25, 26,	A method of making a composition comprising a nucleotide
	27		modification (insertion) to encode a non-PIV molecule where
			the molecule is a <u>cytokine</u> , classified in class 435, subclass 5.
56	1, 21,	25, 28	A method of making a composition comprising a nucleotide
	27		modification (insertion) to encode a non-PIV molecule where
			the molecule is a T-helper epitope , classified in class 435,
			subclass 5.
57	1, 21,	25, 28	A method of making a composition comprising a nucleotide
	27		modification (insertion) to encode a non-PIV molecule where
			the molecule is a <u>restriction site marker</u> , classified in class
			435, subclass 5 <u>.</u>
58	1, 21,	25, 28	A method of making a composition comprising a nucleotide
	27		modification (insertion) to encode a non-PIV molecule where
			the molecule is a protein of a microbial (bacteria or virus)
			pathogen, classified in class 435, subclass 5.
59	1, 27,	32, 33, 35-	A method of making a composition comprising HPIV2
		40, 42	chimera with a heterologous gene encoding antigenic
			determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV1 HN , classified in class 435,
		:	subclass 5.
60	1, 27	32, 33, 37,	A method of making a composition comprising HPIV2
		39	chimera with a heterologous gene encoding antigenic
			determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV1 F , classified in class 435, subclass
			5.
61	1, 27	32, 33, 37,	A method of making a composition comprising HPIV2
		39	chimera with a heterologous gene encoding antigenic

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			determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV2 F , classified in class 435, subclass 5.
62	1, 27	32, 33, 37,	A method of making a composition comprising HPIV2
""	-,	39, 42	chimera with a heterologous gene encoding antigenic
		37, 12	determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV3 HN , classified in class 435,
			subclass 5.
63	1, 27	32, 33, 37,	A method of making a composition comprising HPIV2
03	1,27	39, 42	chimera with a heterologous gene encoding antigenic
		37, 42	determinant. The new gene segment is added (supernumerary)
			in addition to the HPIV2 genes. The supernumerary
			heterologous gene is HPIV3 F , classified in class 435, subclass
			5.
64	1, 27	6, 30	A method of making a composition comprising HPIV2
07	1, 27	0, 50	chimera with at least one different HPIV N, P or L protein.
65	1, 27	6, 58-61	A method of making a composition comprising HPIV2/BPIV3
03	1, 27	0, 38-01	chimera with at least one different BPIV3 N, P or L protein.
66	1, 27	31, 32, 33	A method of making a composition comprising HPIV2
00	1,27	31, 32, 33	chimera with a heterologous gene encoding antigenic
			determinant with an added gene sequence. The supernumerary
			heterologous gene is measles virus HA , classified in class 435,
			subclass 5.
67	1, 27	29	A method of making a composition comprising HPIV2
07	1,27		chimera with a heterologous gene includes a regulatory
			element, classified in class 435, subclass 5.
68	1, 27	31, 34	A method of making a composition comprising HPIV2
	1, 2.	1 2 1, 2 .	chimera with a heterologous measles virus pathogen sequence,
			classified in class 435, subclass 5.
69	1, 27	31, 34	A method of making a composition comprising HPIV2
	-,	,	chimera with a heterologous RSV subgroup A pathogen
			sequence.
70	1, 27	31, 34	A method of making a composition comprising HPIV2
		,	chimera with a heterologous RSV subgroup B virus pathogen
			sequence, classified in class 435, subclass 5.
71	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous mumps virus pathogen sequence,
			classified in class 435, subclass 5.
72	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous human papilloma virus
			pathogen sequence, classified in class 435, subclass 5.
73	1, 27	31, 34	A method of making a composition comprising HPIV2

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			chimera with a heterologous HIV type 1 virus pathogen sequence, classified in class 435, subclass 5.
74	1, 27	31, 34	A method of making a composition comprising HPIV2
	′		chimera with a heterologous HIV type 2 virus pathogen
			sequence, classified in class 435, subclass 5.
75	1, 27	31, 34	A of making a composition comprising HPIV2 chimera with a
			heterologous HSV virus pathogen sequence, classified in class
			435, subclass 5.
76	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous cytomegalovirus pathogen
			sequence, classified in class 435, subclass 5.
77	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous rabies virus pathogen sequence,
			classified in class 435, subclass 5.
78	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous human metapneumovirus
			pathogen sequence, classified in class 435, subclass 5.
79	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous Epstein Barr virus pathogen
			sequence, classified in class 435, subclass 5.
80	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous filovirus pathogen sequence,
			classified in class 435, subclass 5.
81	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous bunyavirus pathogen sequence,
			classified in class 435, subclass 5.
82	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous flavivirus pathogen sequence,
	1 05		classified in class 435, subclass 5.
83	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous alphavirus pathogen sequence,
0.4	1.07	21.24	classified in class 435, subclass 5.
84	1, 27	31, 34	A method of making a composition comprising HPIV2
			chimera with a heterologous influenza virus pathogen
0.5	1 27	12 11 15	sequence, classified in class 435, subclass 5.
85	1, 27	43,44, 45	A method of making a composition comprising HPIV2 chimera with an attenuating mutation HPIV3 JS cp45
			Target at 948 and/or 1566 HPIV2 L protein, classified in class
			435, subclass 5.
86	1, 27	43, 46, 47	A method of making a composition comprising HPIV2
00	1,2/	43, 40, 47	chimera with an attenuating mutation from RSV with a
			substitution at corresponding target position Phe 460 in
			HPIV2 L protein, classified in class 435, subclass 5.
87	1, 27	43, 48, 49	A method of making a composition comprising HPIV2
	1,2/	73, 70, 77	11 mondo of making a composition comprising in 172

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			HPIV2 L protein, classified in class 435, subclass 5.
87	1, 27	43, 48, 49	A method of making a composition comprising HPIV2
			chimera with an additional attenuating mutation from BPIV3.
			Here the target at Ser 1724 in the HPIV2 L protein is a
			substitution, classified in class 435, subclass 5.
88	1, 27	43, 48, 48,	A method of making a composition comprising A of making a
		50	composition comprising HPIV2 chimera with an additional
			attenuating mutation from BPIV3. Here the target at Ser 1724
			in the HPIV2 L protein is a deletion , classified in class 435,
			subclass 5.
89	1, 27,	43, 51, 52,	A method of making a composition comprising HPIV2
		53-56, 62-66	chimera that comprises an additional attenuating mutation
			(insertion deletion, substitution) resulting in any one of the
			following phenotypic changes:
			Attenuation
			Temperature sensitivity
			Cold-adaptation
			Plaque size
			Host-range restriction
			Change in immunogenicity
			Here the attenuation is a partial or complete deletion of HPIV2
			V, classified in class 435, subclass 5.
90		183, 232	A method of vaccinating a subject to elicit an immune response
			against PIV, classified in class 435, subclass 5.

The inventions are distinct, each from the other because of the following reasons:

Inventions (1)-(47) are unrelated. Inventions are unrelated if it can be shown that they are not disclosed as capable of use together and they have different modes of operation, different functions, or different effects (MPEP § 806.04, MPEP § 808.01). In the instant case the different inventions, represent structurally different viral particles having different polynucleotides. Therefore, where structural identity is required, such as for hybridization or expression using the polynucleotide, the different sequences have different effects. The different sequences also will encode different epitopes, so that the protein they encode have different effects.

Groups 1-47 are compositions and are distinct from groups 48-90 which are drawn to methods. Groups 1-47 are compositions and each is distinct from the other because they contain different materials. Though there may be overlap for these groups, the search for one group (structure) will not be coextensive with that of the other group.

Groups 48-89 are drawn to methods and each is distinct from the other because they utilize different starting materials, therefore the outcomes are not be expected to be the same. Groups 48-89 are drawn to a method of producing HPIV2 viral particles having different mutations, growth characteristics or epitopes. Though there may be overlap between these methods in question for groups 48-89, each utilizes different materials and therefore the outcome is expected to be different. Group 90 is a method for immunizing a subject. The method of group 90 uses different steps from the other methods of group 48-89, thereby setting it apart. It differs from the other methods by utilizing different starting materials and techniques, the outcome would therefore not be expected to be the same.

Inventions 48-54 and 1-7 are related as process of making and product made. The inventions are distinct if either or both of the following can be shown: (1) that the process as claimed can be used to make another and materially different product or (2) that the product as claimed can be made by another and materially different process (MPEP § 806.05(f)). In the instant case viruses can be mutated or attenuated by the prolonged incubation of the virus in cell culture or by chemical mutagenesis.

Inventions 1-47 and 90 are related as product and process of use. The inventions can be shown to be distinct if either or both of the following can be shown: (1) the process for using the product as claimed can be practiced with another materially different product or (2) the product as claimed can be used in a materially different process of using that product. See MPEP § 806.05(h). In the instant case a subject can be immunized using isolated and inactivated viral compositions. In the alternative the virus particles can be used to produce proteins that can them be used in detection assays as well as immunization procedures.

Claims 67 link(s) inventions of groups 1-47. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 67. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 67, 89 link(s) inventions of groups 3-47. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 67 and 89. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 67, 70 link(s) inventions of groups 1-2. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 67 and 70. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 67, 73 link(s) inventions of groups 3-6. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 67 and 73. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 67, 83 link(s) inventions of groups 7-10. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 67 and 83. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claim 1 link(s) inventions of groups 48-89. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claim 1. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 1 and 27 link(s) inventions of groups 55-89. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 1 and 27. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the

linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

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Claims 1, 9 link(s) inventions of groups 50-53. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 1 and 9. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Claims 1, 21 link(s) inventions of groups 54-58. The restriction requirement among the linked inventions is subject to the nonallowance of the linking claim(s), claims 1 and 21. Upon the indication of allowability of the linking claim(s), the restriction requirement as to the linked inventions shall be withdrawn and any claim(s) depending from or otherwise requiring all the limitations of the allowable linking claim(s) will be rejoined and fully examined for patentability in accordance with 37 CFR 1.104 Claims that require all the limitations of an allowable linking claim will be entered as a matter of right if the amendment is presented prior to final rejection or allowance, whichever is earlier. Amendments submitted after final rejection are governed by 37 CFR 1.116; amendments submitted after allowance are governed by 37 CFR 1.312.

Applicant(s) are advised that if any claim(s) including all the limitations of the allowable linking claim(s) is/are presented in a continuation or divisional application, the claims of the continuation or divisional application may be subject to provisional statutory and/or nonstatutory double patenting rejections over the claims of the instant application. Where a restriction requirement is withdrawn, the provisions of 35 U.S.C. 121 are no longer applicable. In re Ziegler, 443 F.2d 1211, 1215, 170 USPQ 129, 131-32 (CCPA 1971). See also MPEP § 804.01.

Because these inventions are distinct for the reasons given above, have acquired a separate status in the art as shown by their different classification, and the literature and sequence searches required for each of the Groups are not required for another of the Groups, restriction for examination purposes as indicated is proper.

Applicant is advised that the reply to this requirement to be complete must include an election of the invention to be examined even though the requirement be traversed (37 CFR 1.143).

Applicant is reminded that upon the cancellation of claims to a non-elected invention, the inventorship must be amended in compliance with 37 CFR 1.48(b) if one or more of the currently named inventors is no longer an inventor of at least one claim remaining in the application. Any amendment of inventorship must be accompanied by a petition under 37 CFR 1.48(b) and by the fee required under 37 CFR 1.17(I).

The examiner has required restriction between product and process claims. Where applicant elects claims directed to the product, and the product claims are subsequently found allowable, withdrawn process claims that depend from or otherwise require all the limitations of the allowable product claim will be considered for rejoinder. All claims directed a nonelected process invention must require all the limitations of an allowable product claim for that process invention to be rejoined.

In the event of rejoinder, the requirement for restriction between the product claims and the rejoined process claims will be withdrawn, and the rejoined process claims will be fully examined for patentability in accordance with 37 CFR 1.104. Thus, to be allowable, the rejoined claims must meet all criteria for patentability including the requirements of 35 U.S.C. 101, 102, 103 and 112. Until all claims to the elected product are found allowable, an otherwise proper restriction requirement between product claims and process claims may be maintained. Withdrawn process claims that are not commensurate in scope with an allowable product claim will not be rejoined. See MPEP § 821.04(b). Additionally, in order to retain the right to rejoinder in accordance with the above policy, applicant is advised that the process claims should be amended during prosecution to require the limitations of the product claims. Failure to do so may result in a loss of the right to rejoinder. Further, note that the prohibition against double patenting rejections of 35 U.S.C. 121 does not apply where the restriction requirement is withdrawn by the examiner before the patent issues. See MPEP § 804.01.

Conclusion

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Agnieszka Boesen whose telephone number is 571-272-8035. The examiner can normally be reached on 9:00 AM to 5:30 PM.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Bruce Campell can be reached on 571-272-0974. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Art Unit: 1648

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

AB

Agnieszka Boesen, Ph.D.

Examiner

may 4,2006

ULRIKE WINKLER, PH.D. PRIMARY EXAMINER